



SEQUENCE LISTING

<110> AMYLIN PHARMACEUTICALS, INC

<120> METHODS AND COMPOSITIONS FOR TREATING POLYCYSTIC OVARY SYNDROME

<130> 18528.636

<140> 10/629,649

<141> 2003-07-30

<150> 10/317,126

<151> 2002-12-11

<150> PCT/US03/01109

<151> 2003-01-14

<160> 211

<170> PatentIn version 3.2

<210> 1

<211> 37;

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: Mammalian GLP peptide

<400> 1

His Asp Glu Phe Glu Arg His Ala Glu Gly Thr Phe Thr Ser Asp Val
1 5 10 15

Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu
20 25 30

Val Lys Gly Arg Gly
35

<210> 2

<211> 36

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: Mammalian GLP peptide

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His Asp Glu Phe Glu Arg His Ala Glu Gly Thr Phe Thr Ser Asp Val
1 5 10 15

Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu
20 25 30

Val Lys Gly Arg

<210> 3
 <211> 31
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: Mammalian GLP peptide

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 His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
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 Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg Gly
 20 25 30

<210> 4
 <211> 30
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: Mammalian GLP peptide

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 His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
 1 5 10 15
 Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
 20 25 30

<210> 5
 <211> 29
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: Truncated form of GLP-1

<400> 5
 Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly Gln Ala
 1 5 10 15
 Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg Gly
 20 25

<210> 6
 <211> 28
 <212> PRT
 <213> Unknown

<220>

<223> Description of Unknown Organism: Truncated form of GLP-1

<400> 6

Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly Gln Ala
1 5 10 15

Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
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<210> 7

<211> 39

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: Exendin 3

<400> 7

His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

<210> 8

<211> 31

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: Exendin 4 (9-39) (NH2)

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Asp Leu Ser Lys Gln Met Glu Glu Glu Ala Val Arg Leu Phe Ile Glu
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Trp Leu Lys Asn Gly Gly Pro Ser Ser Gly Ala Pro Pro Pro Ser
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<210> 9

<211> 39

<212> PRT

<213> Unknown

<220>

<223> Description of Unknown Organism: Exendin 4

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His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
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Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
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<210> 10
<211> 38
<212> PRT
<213> Unknown

<220>

<223> Description of Unknown Organism: Helospectin I

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His Ser Asp Ala Thr Phe Thr Ala Glu Tyr Ser Lys Leu Leu Ala Lys
1 5 10 15

Leu Ala Leu Gln Lys Tyr Leu Glu Ser Ile Leu Gly Ser Ser Thr Ser
20 25 30

Pro Arg Pro Pro Ser Ser
35

<210> 11
<211> 37
<212> PRT
<213> Unknown

<220>

<223> Description of Unknown Organism: Helospectin II

<400> 11

His Ser Asp Ala Thr Phe Thr Ala Glu Tyr Ser Lys Leu Leu Ala Lys
1 5 10 15

Leu Ala Leu Gln Lys Tyr Leu Glu Ser Ile Leu Gly Ser Ser Thr Ser
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Pro Arg Pro Pro Ser
35

<210> 12
<211> 35
<212> PRT
<213> Unknown

<220>

<223> Description of Unknown Organism: Helodermin

<400> 12

His Ser Asp Ala Ile Phe Thr Glu Glu Tyr Ser Lys Leu Leu Ala Lys
1 5 10 15

Leu Ala Leu Gln Lys Tyr Leu Ala Ser Ile Leu Gly Ser Arg Thr Ser
 20 25 30

Pro Pro Pro
 35

<210> 13
 <211> 35
 <212> PRT
 <213> Unknown

<220>
 <223> Description of Unknown Organism: Q8, Q9 heliodermin

<400> 13
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 1 5 10 15

Leu Ala Leu Gln Lys Tyr Leu Ala Ser Ile Leu Gly Ser Arg Thr Ser
 20 25 30

Pro Pro Pro
 35

<210> 14
 <211> 30
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 <213> Artificial sequence

<220>
 <223> Truncated form of exendin-4

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 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly
 20 25 30

<210> 15
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 <213> Artificial sequence

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 <223> Truncated form of Exendin-4

<220>
 <221> Variant
 <222> (30)..(30)
 <223> Gly-NH2

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His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Xaa
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<210> 16

<211> 28

<212> PRT

<213> Artificial Sequence

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<223> Truncated form of Exendin-4

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Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 17

<211> 39

<212> PRT

<213> Artificial sequence

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<223> 14-Leu, 25-Phe form of exendin-4

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<221> variant

<222> (39)..(39)

<223> Ser-NH₂

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20 25 30

Ser Gly Ala Pro Pro Pro Xaa
35

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Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Xaa
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<210> 19
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<213> Artificial Sequence

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<223> 14-Leu, 22-Ala, 25-Phe form of exendin-4(1-28)

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<221> variat
<222> (28)..(28)
<223> Asn-NH2

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His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
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Glu Ala Val Arg Leu Ala Ile Glu Phe Leu Lys Xaa
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<223> See specification as filed for detailed description of substitutions and preferred embodiments.

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<222> (2)

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<221> VARIANT

<222> (3)

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 <222> (27)
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 <222> (28)
 <223> Ala or Asn

 <220>
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 <222> (29)..(29)
 <223> -OH, -NH2, Gly-OH, Gly-NH2, Gly Gly-ON, Gly Gly-NH2 and further
 as indicated in the specification

 <400> 20
 Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15

 Xaa Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 20 25

 <210> 21
 <211> 28
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 <223> See specification as filed for detailed description of
 substitutions and preferred embodiments.

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 <222> (2)
 <223> Ser, Gly, Ala, or Thr

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 <222> (3)
 <223> Ala, Asp, or Glu

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<221> VARIANT
 <222> (4)
 <223> Ala, norvaline, Val, norleucine or Gly

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 <222> (5)
 <223> Ala or Thr

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 <222> (9)
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 <222> (10)
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 <222> (11)
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 <222> (13)
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 <222> (14)
 <223> Ala, Leu, Ile, pentylglycine, Val or Met

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<223> Ala or Glu

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 Met

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 <223> Ala, Trp, Phe, Tyr or naphthylalanine

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as indicated in the specification

<400> 21

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1 5 10 15

Xaa Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
20 25

<210> 22

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<223> See specification as filed for detailed description of
substitutions and preferred embodiments.

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<221> VARIANT

<222> (2)..(2)

<223> Gly or Ala

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<221> VARIANT

<222> (3)..(3)

<223> Asp or Glu

<220>

<221> VARIANT

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<223> Ala or Thr

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<221> VARIANT

<222> (6)..(6)

<223> Ala, Phe or naphthalanine

<220>

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<222> (7)..(7)

<223> Thr or Ser

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<222> (8)..(8)

<223> Ala, Ser or Thr

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 <222> (9)..(9)
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 <222> (10)..(10)
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 <223> Ala or Lys

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 <223> Ala or Gln

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 <222> (14)..(14)
 <223> Ala, Leu or pentylglycine

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 <223> Ala or Arg

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 <223> Ile, Val or tert-butylglycine

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 <222> (28)..(28)
 <223> Ala or Asn

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 <223> -OH, -NH2, Gly-OH, Gly-NH2, Gly Gly-ON, Gly Gly-NH2 and further
 as indicated in the specification

 <400> 22

 Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15

 Xaa Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 20 25

 <210> 23
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 <222> (2)..(2)
 <223> Gly or Ala

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 <222> (3)..(3)
 <223> Ala, Asp or Glu

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 <222> (4)..(4)
 <223> Ala or Gly

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 <222> (5)..(5)
 <223> Ala or Thr

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 <222> (9)..(9)
 <223> Ala, Asp or Glu

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 <222> (10)..(10)
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 <222> (13)..(13)
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 <222> (14)..(14)
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 <222> (29)..(29)
 <223> -OH, -NH2, Gly-OH, Gly-NH2, Gly Gly-ON, Gly Gly-NH2 and further
 as indicated in the specification
 <400> 23
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15
 Xaa Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 20 25
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 <223> Ala or Thr

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 or branched alkanoyl or cycloalkanoyl
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 Ala Lys-NH3-R where R is Lys, Arg, C1-C10 straight-chain or
 branched alkanoyl or cycloalkylalkanoyl
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 <222> (28)..(28)
 <223> -OH, -NH2, Gly-OH, Gly-NH2, Gly Gly-ON, Gly Gly-NH2 and further
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Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15

Xaa Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
20 25

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<223> See specification as filed for detailed description of
substitutions and preferred embodiments.

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4imidazopropionyl

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 <222> (10)..(10)
 <223> Ala, Leu, Ile, Val, pentylglycine or Met

<220>
 <221> VARIANT
 <222> (11)..(11)
 <223> Ala or Ser

<220>
 <221> VARIANT
 <222> (12)..(12)
 <223> Ala or Lys

<220>
 <221> VARIANT
 <222> (13)..(13)
 <223> Ala or Gln

<220>
 <221> VARIANT
 <222> (14)..(14)
 <223> Ala, Leu, Ile, pentylglycine, Val or Met

<220>
 <221> VARIANT
 <222> (15)..(17)
 <223> Ala or Glu

<220>
 <221> VARIANT
 <222> (19)..(19)
 <223> Ala or Val

<220>
 <221> VARIANT
 <222> (20)..(20)
 <223> Ala or Arg

<220>
 <221> VARIANT
 <222> (21)..(21)
 <223> Ala, Leu or Lys-NH3-R where R is Lys, Arg, C1-10 straight-chain
 or branched alkanoyl or cycloalylel-alkanoyl

<220>
 <221> VARIANT
 <222> (22)..(22)
 <223> Phe, Tyr or naphthylalanine

<220>
 <221> VARIANT
 <222> (23)..(23)
 <223> Ile, Val, Leu, pentylglycine, tert-butylglycine or Met

 <220>
 <221> VARIANT
 <222> (24)..(24)
 <223> Ala, Glu or Asp

 <220>
 <221> VARIANT
 <222> (25)..(25)
 <223> Ala, Trp, Phe, Tyr or naphthylalanine

 <220>
 <221> VARIANT
 <222> (26)..(26)
 <223> Ala or Leu

 <220>
 <221> VARIANT
 <222> (27)..(27)
 <223> Lys Asn, Asn Lys, Lys-NH3-R Asn, Asn Lys-NH3-R, Lys-NH3-R Ala,
 Ala Lys-NH3-R where R is Lys, Arg, C1-C10 straight-chain or
 branched alkanoyl or cycloalkylalkanoyl

 <220>
 <221> VARIANT
 <222> (28)..(28)
 <223> -OH, -NH2, Gly-OH, Gly-NH2, Gly Gly-ON, Gly Gly-NH2 and further
 as indicated in the specification

 <400> 25

 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 1 5 10 15

 Xaa Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 20 25

 <210> 26
 <211> 39
 <212> PRT
 <213> Artificial sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MISC_FEATURE
 <223> See specification as filed for detailed description of
 substitutions and preferred embodiments.

 <220>

<221> VARIANT
 <222> (1)..(1)
 <223> His, Arg or Tyr

 <220>
 <221> VARIANT
 <222> (2)..(2)
 <223> Ser, Gly, Ala or Thr

 <220>
 <221> VARIANT
 <222> (3)..(3)
 <223> Asp or Glu

 <220>
 <221> VARIANT
 <222> (6)..(6)
 <223> Phe, Tyr or naphthylalanine

 <220>
 <221> VARIANT
 <222> (7)..(7)
 <223> Thr or Ser

 <220>
 <221> VARIANT
 <222> (8)..(8)
 <223> Thr or Ser

 <220>
 <221> VARIANT
 <222> (9)..(9)
 <223> Asp or Glu

 <220>
 <221> VARIANT
 <222> (10)..(10)
 <223> Leu, Ile, Val, pentylglycine or Met

 <220>
 <221> VARIANT
 <222> (14)..(14)
 <223> Leu, Ile, pentylglycine, Val or Met

 <220>
 <221> VARIANT
 <222> (22)..(22)
 <223> Phe, Tyr or naphthylalanine

 <220>
 <221> VARIANT
 <222> (23)..(23)
 <223> Ile, Val, Leu, pentylglycine, tert-butylglycine or Met

 <220>
 <221> VARIANT
 <222> (24)..(24)

<223> Glu or Asp
 <220>
 <221> VARIANT
 <222> (25)..(25)
 <223> Trp, Phe, Tyr or naphthylalanine
 <220>
 <221> VARIANT
 <222> (31)..(31)
 <223> Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine,
 N-alkylpentylglycine or N-alkylalanine
 <220>
 <221> VARIANT
 <222> (36)..(38)
 <223> Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine,
 N-alkylpentylglycine or N-alkylalanine
 <220>
 <221> VARIANT
 <222> (39)..(39)
 <223> Ser, Thr or Tyr and is optionally amidated
 <220>
 <221> VARIANT
 <222> (39)..(39)
 <223> Ser-OH, Ser-NH3, Thr-OH, Thr-NH3, Tyr-OH or Tyr-NH3
 <400> 26
 Xaa Xaa Xaa Gly Thr Xaa Xaa Xaa Xaa Xaa Ser Lys Gln Xaa Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Xaa Xaa Xaa Xaa Leu Lys Asn Gly Gly Xaa Ser
 20 25 30
 Ser Gly Ala Xaa Xaa Xaa Xaa
 35
 <210> 27
 <211> 38
 <212> PRT
 <213> Artificial sequence
 <220>
 <223> artificial sequence with specific variable residues
 <220>
 <221> MISC_FEATURE
 <223> See specification as filed for detailed description of
 substitutions and preferred embodiments.
 <220>

<221> VARIANT
 <222> (1)..(1)
 <223> His, Arg, Tyr or 4-imidazopropionyl

 <220>
 <221> VARIANT
 <222> (2)..(2)
 <223> Ser, Gly, Ala or Thr

 <220>
 <221> VARIANT
 <222> (3)..(3)
 <223> Asp or Glu

 <220>
 <221> VARIANT
 <222> (6)..(6)
 <223> Phe, Tyr or naphthylalanine

 <220>
 <221> VARIANT
 <222> (7)..(7)
 <223> Thr or Ser

 <220>
 <221> VARIANT
 <222> (8)..(8)
 <223> Ser or Thr

 <220>
 <221> VARIANT
 <222> (9)..(9)
 <223> Asp or Glu

 <220>
 <221> VARIANT
 <222> (10)..(10)
 <223> Leu, Ile, Val, pentylglycine or Met

 <220>
 <221> VARIANT
 <222> (14)..(14)
 <223> Leu, Ile, pentylglycine, Val or Met

 <220>
 <221> VARIANT
 <222> (22)..(22)
 <223> Phe, Tyr or naphthylalanine

 <220>
 <221> VARIANT
 <222> (23)..(23)
 <223> Ile, Val, Leu, pentylglycine, tert-butylglycine or Met

 <220>
 <221> VARIANT
 <222> (24)..(24)

<223> Glu or Asp
 <220>
 <221> VARIANT
 <222> (25)..(25)
 <223> Trp, Phe, Tyr or naphthylalanine
 <220>
 <221> VARIANT
 <222> (27)..(27)
 <223> Lys, Asn, Asn, Lys, Lys-NH3-R Asn, Asn, Lys-NH3-R where R is Lys, Arg, C1-C10 straight-chain or branched alkanoyl or cycloalkylalkanoyl
 <220>
 <221> VARIANT
 <222> (30)..(30)
 <223> Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine, N-alkylpentylglycine or N-alkylalanine
 <220>
 <221> misc_feature
 <222> (30)..(30)
 <223> Xaa can be any naturally occurring amino acid
 <220>
 <221> VARIANT
 <222> (35)..(37)
 <223> Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine, N-alkylpentylglycine or N-alkylalanine
 <220>
 <221> VARIANT
 <222> (38)..(38)
 <223> Ser, Thr or Tyr, which is optionally amidated
 <220>
 <221> VARIANT
 <222> (38)..(38)
 <223> Ser-OH, Ser-NH3, Thr-OH, Thr-NH3, Tyr-OH or Tyr-NH3
 <400> 27
 Xaa Xaa Xaa Gly Thr Xaa Xaa Xaa Xaa Xaa Ser Lys Gln Xaa Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Xaa Xaa Xaa Xaa Leu Xaa Gly Gly Xaa Ser Ser
 20 25 30
 Gly Ala Xaa Xaa Xaa Xaa
 35
 <210> 28
 <211> 30

<212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <400> 28
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly
 20 25 30

 <210> 29
 <211> 30
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (30)
 <223> OPTIONAL AMIDATION, Position 30 may be Gly-NH2

 <400> 29
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly
 20 25 30

 <210> 30
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 30
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Ala Ile Glu Phe Leu Lys Asn
 20 25

 <210> 31

<211> 39
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (39)
<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 31
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
20 25 30
Ser Gly Ala Pro Pro Pro Ser
35

<210> 32
<211> 39
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (39)
<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 32
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30
Ser Gly Ala Pro Pro Pro Ser
35

<210> 33
<211> 39
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 33

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

<210> 34

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 34

Tyr Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

<210> 35

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Tyr-NH2

<400> 35

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Tyr
35

<210> 36

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 36

His Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

<210> 37

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<220>

<221> VARIANT

<222> (6)

<223> Xaa is naphthylalanine

<400> 37

His Gly Glu Gly Thr Xaa Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

<210> 38
<211> 39
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (39)
<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 38
His Gly Glu Gly Thr Phe Ser Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30
Ser Gly Ala Pro Pro Pro Ser
35

<210> 39
<211> 39
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (39)
<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 39
His Gly Glu Gly Thr Phe Ser Thr Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30
Ser Gly Ala Pro Pro Pro Ser
35

<210> 40
<211> 39
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

 <400> 40
 His Gly Glu Gly Thr Phe Thr Thr Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

 Ser Gly Ala Pro Pro Pro Ser
 35

<210> 41
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

 <400> 41
 His Gly Glu Gly Thr Phe Thr Ser Glu Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

 Ser Gly Ala Pro Pro Pro Ser
 35

<210> 42
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<220>
 <221> VARIANT
 <222> (10)
 <223> Xaa is pentylglycine

<400> 42
 His Gly Glu Gly Thr Phe Thr Ser Asp Xaa Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30
 Ser Gly Ala Pro Pro Pro Ser
 35

<210> 43
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<220>
 <221> VARIANT
 <222> (10)
 <223> Xaa is pentyglycine

<400> 43
 His Gly Glu Gly Thr Phe Thr Ser Asp Xaa Ser Lys Gln Leu Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30
 Ser Gly Ala Pro Pro Pro Ser
 35

<210> 44
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<220>
 <221> VARIANT
 <222> (14)

<223> Xaa is pentyglycine

<400> 44

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Xaa Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

<210> 45

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<220>

<221> VARIANT

<222> (14)

<223> Xaa is pentyglycine

<400> 45

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Xaa Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

<210> 46

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<220>

<221> VARIANT

<222> (22)
 <223> Xaa is naphthylalanine

<400> 46
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Xaa Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly Ala Pro Pro Pro Ser
 35

<210> 47
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 47
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Val Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly Ala Pro Pro Pro Ser
 35

<210> 48
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 48
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Val Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

<210> 49

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (23)

<223> Xaa at Position 23 is tertiary-butylglycine

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 49

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Xaa Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

<210> 50

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (23)

<223> Xaa at position 23 is tertiary-butylglycine

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 50

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Xaa Glu Phe Leu Lys Asn Gly Gly Pro Ser

30

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (31)

<223> Xaa at position 31 is thioproline

<220>

<221> VARIANT

<222> (36)..(38)

<223> Xaa at positions 36, 36, and 38 is thioproline

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 53

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Met	Glu	Glu
1				5					10					15	

Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Trp	Leu	Lys	Asn	Gly	Gly	Xaa	Ser
			20					25					30		

Ser	Gly	Ala	Xaa	Xaa	Xaa	Ser
			35			

<210> 54

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (36)..(38)

<223> Xaa at positions 36, 37, and 38 is thioproline

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 54

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Met	Glu	Glu
1				5					10					15	

Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Trp	Leu	Lys	Asn	Gly	Gly	Pro	Ser
			20					25					30		

Ser	Gly	Ala	Xaa	Xaa	Xaa	Ser
-----	-----	-----	-----	-----	-----	-----

<210> 55
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (31)
 <223> Xaa at position 31 is homoproline

<220>
 <221> VARIANT
 <222> (36)..(38)
 <223> Xaa at positions 36, 37, and 38 is homoproline

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 55
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
 20 25 30

Ser Gly Ala Xaa Xaa Xaa Ser
 35

<210> 56
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (36)..(38)
 <223> Xaa at positions 36, 37, and 38 is homoproline

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 56
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu

1	5	10	15												
Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Trp	Leu	Lys	Asn	Gly	Gly	Pro	Ser
			20					25					30		

Ser Gly Ala Xaa Xaa Xaa Ser
35

<210> 57
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (31)
 <223> Xaa at position 31 is thioproline

<220>
 <221> VARIANT
 <222> (36)..(38)
 <223> Xaa at positions 36, 37, and 38 is thioproline

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 57
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Phe	Leu	Lys	Asn	Gly	Gly	Xaa	Ser
			20					25					30		

Ser Gly Ala Xaa Xaa Xaa Ser
35

<210> 58
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (31)
 <223> Xaa at position 31 is homoproline

<220>

<221> VARIANT
 <222> (36)..(38)
 <223> Xaa at positions 36,37, and 38 is homoproline

 <220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

 <400> 58
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Xaa Ser
 20 25 30

 Ser Gly Ala Xaa Xaa Xaa Ser
 35

<210> 59
 <211> 39
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (31)
 <223> Xaa at position 31 is N-methylalanine

<220>
 <221> VARIANT
 <222> (36)..(38)
 <223> Xaa at positions 36, 37, and 38 is N-methylalanine

<220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 59
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
 20 25 30

 Ser Gly Ala Xaa Xaa Xaa Ser
 35

<210> 60
 <211> 39
 <212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (36)..(38)

<223> Xaa at positions 36, 37, and 38 is N-methylalanine

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 60

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Met	Glu	Glu
1				5					10					15	

Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Trp	Leu	Lys	Asn	Gly	Gly	Pro	Ser
			20					25					30		

Ser Gly Ala Xaa Xaa Xaa Ser

35

<210> 61

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (31)

<223> Xaa at position 31 is N-methylalanine

<220>

<221> VARIANT

<222> (36)..(38)

<223> Xaa at positions 36, 37, and 38 is N-methylalanine

<220>

<221> MOD_RES

<222> (39)

<223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

<400> 61

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Leu	Glu	Glu
1				5					10					15	

Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Phe	Leu	Lys	Asn	Gly	Gly	Xaa	Ser
			20					25					30		

Ser Gly Ala Xaa Xaa Xaa Ser

<210> 62
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 62
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

<210> 63
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 63
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 64
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 64
 His Ala Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 65
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 65
 His Gly Glu Gly Ala Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 66
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
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<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 66
 His Gly Glu Gly Thr Ala Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 67
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 67

His Gly Glu Gly Thr Phe Thr Ala Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 68

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 68

His Gly Glu Gly Thr Phe Thr Ser Asp Ala Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 69

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 69

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ala Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 70
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 70
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Ala Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

 <210> 71
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 71
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Ala Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

 <210> 72
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 72
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Ala Glu Glu

```

1              5              10              15
Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
      20              25

<210> 73
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 73
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Ala Glu
1              5              10              15
Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
      20              25

<210> 74
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 74
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Ala
1              5              10              15
Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
      20              25

<210> 75
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>

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<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 75
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1          5          10          15
Ala Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
          20          25

<210> 76
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 76
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1          5          10          15
Glu Ala Ala Arg Leu Phe Ile Glu Phe Leu Lys Asn
          20          25

<210> 77
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 77
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1          5          10          15
Glu Ala Val Ala Leu Phe Ile Glu Phe Leu Lys Asn
          20          25

<210> 78
<211> 28
<212> PRT

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<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 78

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Leu	Glu	Glu
1				5					10				15		

Glu	Ala	Val	Arg	Ala	Phe	Ile	Glu	Phe	Leu	Lys	Asn
			20				25				

<210> 79

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 79

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Leu	Glu	Glu
1				5					10				15		

Glu	Ala	Val	Arg	Leu	Phe	Ile	Ala	Phe	Leu	Lys	Asn
			20				25				

<210> 80

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 80

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Leu	Glu	Glu
1				5					10				15		

Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Ala	Leu	Lys	Asn
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

<210> 81
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 81
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Phe Ala Lys Asn
 20 25

<210> 82
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 82
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Ala Asn
 20 25

<210> 83
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Ala-NH2

<400> 83
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Ala
 20 25

<210> 84
 <211> 38
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (38)
 <223> OPTIONAL AMIDATION, Position 38 may be Pro-NH2

<400> 84
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly Ala Pro Pro Pro
 35

<210> 85
 <211> 38
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (38)
 <223> OPTIONAL AMIDATION, Position 38 may be Pro-NH2

<400> 85
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly Ala Pro Pro Pro
 35

<210> 86
<211> 37
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (37)
<223> OPTIONAL AMIDATION, Position 37 may be Pro-NH2

<400> 86
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly Ala Pro Pro
 35

<210> 87
<211> 37
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (37)
<223> OPTIONAL AMIDATION, Position 37 may be Pro-NH2

<400> 87
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly Ala Pro Pro
 35

<210> 88
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES
 <222> (36)
 <223> OPTIONAL AMIDATION, Position 36 may be Pro-NH2

 <400> 88
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30
 Ser Gly Ala Pro
 35

<210> 89
 <211> 36
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (36)
 <223> OPTIONAL AMIDATION, Position 36 may be Pro-NH2

 <400> 89
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30
 Ser Gly Ala Pro
 35

<210> 90
 <211> 35
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (35)
 <223> OPTIONAL AMIDATION, Position 35 may be Ala-NH2

 <400> 90
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser

20

25

30

Ser Gly Ala
35

<210> 91

<211> 35

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (35)

<223> OPTIONAL AMIDATION, Position 35 may be Ala-NH2

<400> 91

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala
35

<210> 92

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (34)

<223> OPTIONAL AMIDATION, Position 34 may be Gly-NH2

<400> 92

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly

<210> 93

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (34)

<223> OPTIONAL AMIDATION, Position 34 may be Gly-NH2

<400> 93

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly

<210> 94

<211> 33

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (33)

<223> OPTIONAL AMIDATION, Position 33 may be Ser-NH2

<400> 94

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser

<210> 95

<211> 33

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (33)

<223> OPTIONAL AMIDATION, Position 33 may be Ser-NH2

<400> 95

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser

<210> 96

<211> 32

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (32)

<223> OPTIONAL AMIDATION, Position 32 may be Ser-NH2

<400> 96

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

<210> 97

<211> 32

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (32)

<223> OPTIONAL AMIDATION, Position 32 may be Ser-NH2

<400> 97

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
20 25 30

<210> 98

<211> 31

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (31)

<223> OPTIONAL AMIDATION, Position 31 may be Pro-NH2

<400> 98

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Met	Glu	Glu
1				5					10					15	

Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Trp	Leu	Lys	Asn	Gly	Gly	Pro
			20					25					30	

<210> 99

<211> 31

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (31)

<223> OPTIONAL AMIDATION, Position 31 may be Pro-NH2

<400> 99

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Leu	Glu	Glu
1				5					10					15	

Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Phe	Leu	Lys	Asn	Gly	Gly	Pro
			20					25					30	

<210> 100

<211> 30

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (30)

<223> OPTIONAL AMIDATION, Position 30 may be Gly-NH2

<400> 100

His	Gly	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Leu	Ser	Lys	Gln	Leu	Glu	Glu
1				5					10					15	

Glu	Ala	Val	Arg	Leu	Phe	Ile	Glu	Phe	Leu	Lys	Asn	Gly	Gly
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

20

25

30

<210> 101
 <211> 29
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (29)
 <223> OPTIONAL AMIDATION, Position 29 may be Gly-NH2

<400> 101
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly
 20 25

<210> 102
 <211> 29
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (29)
 <223> OPTIONAL AMIDATION, Position 29 may be Gly-NH2

<400> 102
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly
 20 25

<210> 103
 <211> 38
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (31)
 <223> Xaa is thioproline

<220>
<221> VARIANT
<222> (36)..(38)
<223> Xaa is thioproline

<220>
<221> MOD_RES
<222> (38)
<223> OPTIONAL AMIDATION, Position 38 may be thioproline-NH2

<400> 103
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
20 25 30
Ser Gly Ala Xaa Xaa Xaa
35

<210> 104
<211> 38
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> VARIANT
<222> (36)..(38)
<223> Xaa is thioproline

<220>
<221> MOD_RES
<222> (38)
<223> OPTIONAL AMIDATION, Position 38 may be thioproline-NH2

<400> 104
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30
Ser Gly Ala Xaa Xaa Xaa
35

<210> 105
<211> 37
<212> PRT
<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (31)

<223> Xaa is N-methylalanine

<220>

<221> MOD_RES

<222> (37)

<223> OPTIONAL AMIDATION, Position 37 may be Pro-NH2

<400> 105

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
20 25 30

Ser Gly Ala Pro Pro
35

<210> 106

<211> 37

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (31)

<223> Xaa is N-methylalanine

<220>

<221> VARIANT

<222> (36)..(37)

<223> Xaa is N-methylalanine

<220>

<221> MOD_RES

<222> (37)

<223> OPTIONAL AMIDATION, Position 37 may be N-methylalanine-NH2

<400> 106

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
20 25 30

Ser Gly Ala Xaa Xaa
35

<210> 107
 <211> 37
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (31)
 <223> Xaa is homoproline

<220>
 <221> VARIANT
 <222> (36)..(37)
 <223> Xaa is homoproline

<220>
 <221> MOD_RES
 <222> (37)
 <223> OPTIONAL AMIDATION, Position 37 may be homoproline-NH2

<400> 107
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
 20 25 30
 Ser Gly Ala Xaa Xaa
 35

<210> 108
 <211> 36
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (31)
 <223> Xaa is homoproline

<220>
 <221> VARIANT
 <222> (36)
 <223> Xaa is homoproline

<220>
 <221> MOD_RES
 <222> (36)
 <223> OPTIONAL AMIDATION, Position 36 may be homoproline-NH2

<400> 108
 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
 20 25 30
 Ser Gly Ala Xaa
 35

<210> 109
 <211> 35
 <212> PRT
 <213> Artificial Sequence

<220>
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<220>
 <221> MOD_RES
 <222> (35)
 <223> OPTIONAL AMIDATION, Position 35 may be Ala-NH2

<400> 109
 Arg Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30
 Ser Gly Ala
 35

<210> 110
 <211> 30
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (30)
 <223> OPTIONAL AMIDATION, Position 30 may be Gly-NH2

<400> 110
 His Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly
 20 25 30

<210> 111

<211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
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 <220>
 <221> VARIANT
 <222> (6)
 <223> Xaa is naphthylalanine

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 111
 His Gly Glu Gly Thr Xaa Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

 <210> 112
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
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 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 112
 His Gly Glu Gly Thr Phe Ser Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

 <210> 113
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
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 <220>
 <221> MOD_RES
 <222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 113

His Gly Glu Gly Thr Phe Ser Thr Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 114

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 114

His Gly Glu Gly Thr Phe Thr Ser Glu Leu Ser Lys Gln Met Ala Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 115

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (10)

<223> Xaa is pentylglycine

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 115

His Gly Glu Gly Thr Phe Thr Ser Asp Xaa Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 116
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
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<220>
<221> VARIANT
<222> (22)
<223> Xaa is naphthylalanine

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 116
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Xaa Ile Glu Phe Leu Lys Asn
20 25

<210> 117
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
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<220>
<221> VARIANT
<222> (23)
<223> Xaa is tertiary-butylglycine

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 117
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Xaa Glu Trp Leu Lys Asn
20 25

<210> 118
<211> 28
<212> PRT
<213> Artificial Sequence

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<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 118
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1          5          10          15
Glu Ala Val Arg Leu Phe Ile Asp Phe Leu Lys Asn
          20          25

<210> 119
<211> 33
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (33)
<223> OPTIONAL AMIDATION, Position 33 may be Ser-NH2

<400> 119
His Gly Glu Gly Thr Phe Thr Ser Asp Ala Ser Lys Gln Leu Glu Glu
1          5          10          15
Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
          20          25          30
Ser

<210> 120
<211> 29
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (29)
<223> OPTIONAL AMIDATION, Position 29 may be Gly-NH2

<400> 120
His Gly Glu Gly Thr Phe Thr Ser Asp Ala Ser Lys Gln Met Glu Glu
1          5          10          15

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Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly
20 25

<210> 121
<211> 37
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> VARIANT
<222> (31)
<223> Xaa is homoproline

<220>
<221> VARIANT
<222> (36)..(37)
<223> Xaa is homoproline

<220>
<221> MOD_RES
<222> (37)
<223> OPTIONAL AMIDATION, Position 37 may be homoproline-NH2

<400> 121
His Gly Glu Gly Thr Phe Thr Ser Asp Ala Ser Lys Gln Met Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
20 25 30

Ser Gly Ala Xaa Xaa
35

<210> 122
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 122
Ala Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 123
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 123
His Gly Ala Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 124
<211> 28
<212> PRT
<213> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 124
His Gly Glu Ala Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 125
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 125
His Gly Glu Gly Thr Phe Thr Ser Ala Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 126
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 126
Ala Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 127
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 127
His Gly Ala Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 128
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES

<220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 131
 Ala Ala Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

 <210> 132
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 132
 Ala Ala Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

 <210> 133
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 133
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

<210> 134
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 134
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 135
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 135
Ala Gly Asp Gly Ala Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 136
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 136

Ala Gly Asp Gly Ala Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 137

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (6)

<223> Xaa is naphthylalanine

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 137

Ala Gly Asp Gly Thr Xaa Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 138

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (6)

<223> Xaa is naphthylalanine

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 138

Ala Gly Asp Gly Thr Xaa Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn

<210> 139
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 139
 Ala Gly Asp Gly Thr Phe Ser Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

<210> 140
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 140
 Ala Gly Asp Gly Thr Phe Ser Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 141
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 141
 Ala Gly Asp Gly Thr Phe Thr Ala Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

<210> 142
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 142
 Ala Gly Asp Gly Thr Phe Thr Ala Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 143
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 143
 Ala Gly Asp Gly Thr Phe Thr Ser Ala Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

<210> 144
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 144

Ala Gly Asp Gly Thr Phe Thr Ser Ala Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 145

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 145

Ala Gly Asp Gly Thr Phe Thr Ser Glu Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 146

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 146

Ala Gly Asp Gly Thr Phe Thr Ser Glu Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 147
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 147
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Ala Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

 <210> 148
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 148
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Ala Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

 <210> 149
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> VARIANT
 <222> (10)
 <223> Xaa is pentylglycine

 <220>
 <221> MOD_RES

<222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 149
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Xaa Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

<210> 150
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (10)
 <223> Xaa is pentylglycine

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 150
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Xaa Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 151
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 151
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ala Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

<210> 152
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 152
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ala Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

 <210> 153
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 153
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Ala Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

 <210> 154
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 154

Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Ala Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 155
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 155
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Ala Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 156
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 156
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Ala Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 157
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 157
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Ala Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

<210> 158
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 158
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Ala Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 159
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (14)
 <223> Xaa is pentylglycine

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 159
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Xaa Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn

<210> 160
 <211> 28
 <212> PRT
 <213> Artificial Sequence ,

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> VARIANT
 <222> (14)
 <223> Xaa is pentylglycine

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 160
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Xaa Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

<210> 161
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 161
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Ala Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
 20 25

<210> 162
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

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<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 162
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Ala Glu
1          5          10          15
Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
          20          25

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<210> 163
<211> 28
<212> PRT
<213> Artificial Sequence
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<220>
<223> artificial sequence with specific variable residues
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<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 163
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Ala
1          5          10          15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
          20          25

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<210> 164
<211> 28
<212> PRT
<213> Artificial Sequence
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<220>
<223> artificial sequence with specific variable residues
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<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2
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<400> 164
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Ala
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 165

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<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 165
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1          5          10         15

Ala Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
          20         25

<210> 166
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 166
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1          5          10         15

Ala Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
          20         25

<210> 167
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 167
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1          5          10         15

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Glu Ala Ala Arg Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 168
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 168
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Ala Arg Leu Phe Ile Glu Phe Leu Lys Asn
20 25

<210> 169
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 169
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Ala Leu Phe Ile Glu Trp Leu Lys Asn
20 25

<210> 170
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES

<222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 170
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Ala Leu Phe Ile Glu Phe Leu Lys Asn
 20 25

 <210> 171
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 171
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Ala Phe Ile Glu Trp Leu Lys Asn
 20 25

 <210> 172
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 172
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Ala Phe Ile Glu Phe Leu Lys Asn
 20 25

 <210> 173
 <211> 28
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> VARIANT
 <222> (22)
 <223> Xaa is naphthylalanine

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 173
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Xaa Ile Glu Trp Leu Lys Asn
 20 25

 <210> 174
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> VARIANT
 <222> (22)
 <223> Xaa is naphthylalanine

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 174
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Xaa Ile Glu Phe Leu Lys Asn
 20 25

 <210> 175
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>

<221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 175
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Val Glu Trp Leu Lys Asn
 20 25

 <210> 176
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 176
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Val Glu Phe Leu Lys Asn
 20 25

 <210> 177
 <211> 28
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> VARIANT
 <222> (23)
 <223> Xaa is tertiary-butylglycine

 <220>
 <221> MOD_RES
 <222> (28)
 <223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

 <400> 177
 Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Xaa Glu Trp Leu Lys Asn
 20 25

<210> 178
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> VARIANT
<222> (23)
<223> Xaa is tertiary-butylglycine

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 178
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Xaa Glu Phe Leu Lys Asn
20 25

<210> 179
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 179
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Asp Trp Leu Lys Asn
20 25

<210> 180
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues


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<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 183
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1          5          10         15

Glu Ala Val Arg Leu Phe Ile Glu Trp Ala Lys Asn
          20         25

<210> 184
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 184
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1          5          10         15

Glu Ala Val Arg Leu Phe Ile Glu Phe Ala Lys Asn
          20         25

<210> 185
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 185
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1          5          10         15

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Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Ala Asn
20 25

<210> 186
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Asn-NH2

<400> 186
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Ala Asn
20 25

<210> 187
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)
<223> OPTIONAL AMIDATION, Position 28 may be Ala-NH2

<400> 187
Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Ala
20 25

<210> 188
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (28)

<223> OPTIONAL AMIDATION, Position 28 may be Ala-NH2

<400> 188

Ala Gly Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Ala
20 25

<210> 189

<211> 38

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (38)

<223> OPTIONAL AMIDATION, Position 38 may be Pro-NH2

<400> 189

Ala Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro
35

<210> 190

<211> 38

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> MOD_RES

<222> (38)

<223> OPTIONAL AMIDATION, Position 38 may be Pro-NH2

<400> 190

His Gly Ala Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro
35

<210> 191
<211> 37
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (37)
<223> OPTIONAL AMIDATION, Position 37 may be Pro-NH2

<400> 191
His Gly Glu Ala Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30
Ser Gly Ala Pro Pro
35

<210> 192
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (36)
<223> OPTIONAL AMIDATION, Position 36 may be Pro-NH2

<400> 192
His Gly Glu Gly Thr Phe Thr Ser Ala Leu Ser Lys Gln Met Glu Glu
1 5 10 15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30
Ser Gly Ala Pro
35

<210> 193
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (36)
 <223> OPTIONAL AMIDATION, Position 36 may be Pro-NH2

 <400> 193
 Ala Gly Glu Gly Thr Phe Thr Ser Asp Ala Ser Lys Gln Leu Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30
 Ser Gly Ala Pro
 35

<210> 194
 <211> 35
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (35)
 <223> OPTIONAL AMIDATION, Position 35 may be Ala-NH2

 <400> 194
 Ala Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30
 Ser Gly Ala
 35

<210> 195
 <211> 35
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (35)
 <223> OPTIONAL AMIDATION, Position 35 may be Ala-NH2

<400> 195
 His Gly Ala Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly Ala
 35

<210> 196
 <211> 34
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (34)
 <223> OPTIONAL AMIDATION, Position 34 may be Gly-NH2

<400> 196
 His Gly Glu Ala Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly

<210> 197
 <211> 33
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (33)
 <223> OPTIONAL AMIDATION, Position 33 may be Ser-NH2

<400> 197
 His Gly Glu Gly Thr Phe Thr Ser Ala Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser

<210> 198
 <211> 32

<212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (32)
 <223> OPTIONAL AMIDATION, Position 32 may be Ser-NH2

 <400> 198
 Ala Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

 <210> 199
 <211> 32
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (32)
 <223> OPTIONAL AMIDATION, Position 32 may be Ser-NH2

 <400> 199
 His Gly Ala Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30

 <210> 200
 <211> 31
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (31)
 <223> OPTIONAL AMIDATION, Position 31 may be Pro-NH2

 <400> 200
 His Gly Glu Ala Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro
 20 25 30

<210> 201
 <211> 30
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (30)
 <223> OPTIONAL AMIDATION, Position 30 may be Gly-NH2

<400> 201
 His Gly Glu Gly Thr Phe Thr Ser Ala Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly
 20 25 30

<210> 202
 <211> 29
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> MOD_RES
 <222> (29)
 <223> OPTIONAL AMIDATION, Position 29 may be Gly-NH2

<400> 202
 Ala Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly
 20 25

<210> 203
 <211> 38
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

<220>
 <221> VARIANT
 <222> (31)

<223> Xaa is thioproline

<220>

<221> VARIANT

<222> (36)..(38)

<223> Xaa is thioproline

<220>

<221> MOD_RES

<222> (38)

<223> OPTIONAL AMIDATION, Position 38 may be thioproline-NH2

<400> 203

His Gly Ala Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
 20 25 30

Ser Gly Ala Xaa Xaa Xaa
 35

<210> 204

<211> 38

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (36)..(38)

<223> Xaa is thioproline

<220>

<221> MOD_RES

<222> (38)

<223> OPTIONAL AMIDATION, Position 38 may be thioproline-NH2

<400> 204

His Gly Glu Ala Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30

Ser Gly Ala Xaa Xaa Xaa
 35

<210> 205

<211> 37

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (31)

<223> Xaa is N-methylalanine

<220>

<221> VARIANT

<222> (36)..(37)

<223> Xaa is N-methylalanine

<220>

<221> MOD_RES

<222> (37)

<223> OPTIONAL AMIDATION, Position 37 may be N-methylalanine-NH2

<400> 205

His Gly Glu Gly Thr Phe Thr Ser Ala Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
20 25 30

Ser Gly Ala Xaa Xaa
35

<210> 206

<211> 36

<212> PRT

<213> Artificial Sequence

<220>

<223> artificial sequence with specific variable residues

<220>

<221> VARIANT

<222> (31)

<223> Xaa is homoproline

<220>

<221> VARIANT

<222> (36)

<223> Xaa is homoproline

<220>

<221> MOD_RES

<222> (36)

<223> OPTIONAL AMIDATION, Position 36 may be homoproline-NH2

<400> 206

Ala Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa Ser
20 25 30

Ser Gly Ala Xaa
35

<210> 207
<211> 35
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (35)
<223> OPTIONAL AMIDATION, Position 35 may be Ala-NH2

<400> 207
His Gly Ala Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala
35

<210> 208
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> artificial sequence with specific variable residues

<220>
<221> MOD_RES
<222> (30)
<223> OPTIONAL AMIDATION, Position 30 may be Gly-NH2

<400> 208
His Gly Asp Ala Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly
20 25 30

<210> 209
<211> 39
<212> PRT
<213> Artificial Sequence

<220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

 <400> 209
 Ala Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 20 25 30
 Ser Gly Ala Pro Pro Pro Ser
 35

 <210> 210
 <211> 39
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable residues

 <220>
 <221> MOD_RES
 <222> (39)
 <223> OPTIONAL AMIDATION, Position 39 may be Ser-NH2

 <400> 210
 Ala Gly Ala Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
 1 5 10 15
 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
 20 25 30
 Ser Gly Ala Pro Pro Pro Ser
 35

 <210> 211
 <211> 10
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> artificial sequence with specific variable
 residues

 <220>
 <221> MOD_RES
 <222> 1
 <223> OPTIONAL AMIDATION, Gly at position 1 is optionally

amidated in the case that residues in positions
2-10 are absent

<220>

<221> MOD_RES

<222> 2

<223> OPTIONAL AMIDATION, Gly at position 2 may be optional and
optionally amidated in the case that residues in
positions 3-10 are absent

<220>

<221> VARIANT

<222> 3

<223> Xaa is selected from Pro, homoproline, 3Hyp, 4Hyp,
thioprolin, N-Alkylglycine, N-alkylpentylglycine,
or N-alkylalanine and is optionally amidated in the
case that residues in positions 4...10 are absent

<220>

<221> MOD_RES

<222> 4

<223> OPTIONAL AMIDATION, Ser at position 4 is optionally
amidated in the case that residues in positions
5-10 are absent

<220>

<221> MOD_RES

<222> 5

<223> OPTIONAL AMIDATION, Ser at position 5 is optionally
amidated in the case that residues in positions
6-10 are absent

<220>

<221> MOD_RES

<222> 6

<223> OPTIONAL AMIDATION, Gly at position 6 is optionally
amidated in the case that residues in position
7-10 are absent

<220>

<221> MOD_RES

<222> 7

<223> OPTIONAL AMIDATION, Ala at position 7 is optionally
amidated in the case that residues in positions
8-10 are absent

<220>

<221> VARIANT

<222> 8

<223> Xaa is selected from Pro, homoproline, 3Hyp, 4Hyp,
thioprolin, N-alkylglycine, N-alkylpentylglycine,
or N-alkylalanine and is optionally amidated in
the case that residues in positions 9-10 are absent

<220>

<221> VARIANT

<222> 9

<223> Xaa is selected from Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine, N-alkylpentylglycine, or N-alkylalanine and is optionally amidated in the case that residues in position 10 are absent

<220>

<221> VARIANT

<222> 10

<223> Xaa is selected from Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine, N-alkylpentylglycine, or N-alkylalanine and is optionally amidated

<400> 211

Gly	Gly	Xaa	Ser	Ser	Gly	Ala	Xaa	Xaa	Xaa
1				5				10	